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polyA_site	61 332	?
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gene CDS	1 152	. B 3
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Ç.	Alignment Scores: Pred. No.: Score: 1.57e-51 Matches: 194 Percent Similarity: 100.00\$ Best Local Similarity: 100.00\$ Query Match: 100.00\$ Indels: 0 Gaps: 0 Gaps: 0	. COBBOD
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JOURNAL INE MEDLINE 962 PUBMED 869 REFERENCE 2 AUTHORS Cox	/translation; "MOSHALLIALVAAGLASAQLEDIPPCALINGFVEALGNDGCTRLT DFKCHCSKPELPGQITPCVEEACPLDARISVSHIVVDOCSKAGVPIDIPPVDTTAAPE PSETARETARETEEEPTAEPTAEPTAEPTHEEPTEEPTAVPTGTGGGVPTGTGSFTVTGR PTASTPAEFPGAGSNVRASVGGIAAALLGLAAYL" BASE COUNT 249 a 369 C 275 G 700 L	Œ
REFERENCE 1 AUTHORS Zhu TITLE MOI	/codon_start=1 /product="immunoreactive spherule cell wall protein" /protein_id="AAC49140.1" /dh_vas===cr.151314.1"	
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LOCUS CIL	University of Arizona, 1501 N. Campbell, Tucson, AZ 85724, USA source 11201	
RESULT 2	JOURNAL Submitted (01-NOV) 1995) Frie Committed to the Committed C	
Db 692 (2 (báses 1	-
	JOURNAL Biochem. Biophys. Res. Commun. 218 (2), 485-489 (1996) PIBMED 05:144702	
	Cloning and sequence analysis of the cDNA for a protein f	
Oy 141	EUKBTYOCE; FUngi; Ascomycota; Pezizomycotina; Eurotiomycetes; Onygenales; mitosporic Onygenales; Coccidioides. 1 (bases 1 to 1201)	_
Db 512	SURCE Coccidioides posadasii ORGANISM Coccidioides posadasii	
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Db 452	Complete cds.	
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hu,Y., Yang,C., Magee,D.M. and Cox,R.A.
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nfect. Immun. 64 (7), 2695-2699 (1996) occidioides posadasii
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ukaryota; Pungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
mygenales; mitosporic Onygenales; Coccidioides.

(bases 1 to 1234) rect Submission bmitted (27-JUL-1995) Texas Center for Infectious Disease, search Immunology, 2303/9.E. Military Drive, San Antonio, 1032518 1234 bp mRNA linear PLN 04-NOV-1996 occidioidee immitis antigen 2 (Ag2)-mRNA, complete cde. 32518 GlyGlyIleAlaAlaAlaLeuLeuGlyLeuAlaAlaTyrLeu 194 ProThrGlyThrGlyGlyGlyValProThrGlyThrGlySerPheThrValThrGlyArg 160 Peb 24, 1996 this sequence version replaced gi:1184058. Location/Qualifiers ccaactoccrecaceccaacroaagrreceaagerocregereciático ProThrAlaSerThrProAlaGluPheProGlyAlaGlySerAsnvalArgAlaSerVal 180 ProGluProSerGluThrAlaGluProThrAlaGluProThrGluGluProThrAlaGlu 120 ccinecacrandeceacescraadecaacrantatatatateceateatatatateceateceate 571 ProThrAlaGluProThrAlaGluProThrH18GluProThrGluGluProThrAlaVal 140 bases 1 to 1234) /gene='Ag2'
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